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BLASTP ALIGNMENT OF C-TYPE LECTIN RECEPTOR-LIKE WITH MOUSE MACROPHAGE C-TYPE LECTIN

Score = 412 (145.0 bits), Expect = 1.2e-37, P = 1.2e-37

Identities = 85/215 (39%), Positives = 119/215 (55%), Frame = +1

EEPQDREKGLWWFQLKVWSMAVVSILLLSVCFTVSSVVPHNFMYSKTVKRLSKLREYQQY 234 55

Z ⋳ ++H A++ CF ĽS AVVSI δΓ EE Q + KG

C-Type Lectin Receptor-Like:

SEQ ID NO.

Macrophage C-Type Lectin:

09 EESQMKSKGTRHPQLIPCVFAVVSISFLSACFISTCLVTHHYFLRWTRGSVVKLSDY---

4

235 HSSLTCVME----GKDIEDWSCCPTPWTSFQSSCYFISTGMQSWTKSQKNCSVMGADLVV 402 SEQ ID NO. 4

61 HTRVTCIREEPQPGATGGTWTCCPVSWRAFQSNCYFPLNDNQTWHESERNCSGMSSHLVT 120 Macrophage C-Type Lectin:

W+CCP W +FQS+CYF

G

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H+ +TC+

C-Type Lectin Receptor-Like:

Q+W +S++NCS M + LV

403 INTTEEHDFIIHNLKRNSSYFLGLSHPRGRRHWQWVDHTPYNENVTFWHSGEPNN-LDER 579 C-Type Lectin Receptor-Like: SEQ ID NO.

121 INTEAEQNEVTQLLDKRFSYFLGLADENVEGQWQWVDKTPFNPHTVFWEKGESNDFMEED 180 E +F+ INI

SYFIGI+

+ 1

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+

WQWVD TP+N

580 CAIINERSSQEWGWNDIHCHVPHKSICEMKKIYIYMKYS 696 SEQ ID NO. 4

S × + IC++ CH H+W WND C ++

C-Type Lectin Receptor-Like:

Macrophage C-Type Lectin:

181 CVVL-VHVHEKWVWNDFPCHFEVRRICKLPGITFNWKPS 218 Macrophage C-Type Lectin:

tions their first first first first their times the first first the first times to the first times times to the first times to the first times times times to the first times time

BLASTP ALIGNMENT OF C-TYPE LECTIN RECEPTOR-LIKE WITH DENDRITIC CELL IMMUNORECEPTOR

Score = 529 (186.2 bits), Expect = 4.8e-50, P = 4.8e-50

Identities = 93/188 (498), Positives = 130/188 (698), Frame = +1

118 VVSILLLSVCFTVSSVVPHNFMYSKTVKRLSKLREYQQYHSSLTCVMEGKDIED--WSCC 291

H. LLL++ F ++ V+ + K + L K + L++L CV + +E+ WSCC

C-Type Lectin Receptor-Like:

SEQ ID NO.

51 LIFFLLLAISFFIAFVI----FFQKYSQLLEKKTTKELVHTTLECVKKNMPVEETAWSCC 106 Dendritic Cell Immunoreceptor:

292 PTPWTSFQSSCYFISTGMQSWTKSQKNCSVMGADLVVINTTEEHDFIIHNLKRNSSYFLG 471 S+YF+G NI+ S+K+C+ M A L+VINT EE DFI SW SF S+CYFIST S Д C-Type Lectin Receptor-Like: SEQ ID NO.

Dendritic Cell Immunoreceptor: 107 PKNWKSFSSNCYFISTESASWQDSEKDCARMEAHLLVINTQEEQDFIFQNLQEESAYFVG 166

472 LSHPRGRRHWQWVDHTPYNENVTFWHSGEPNNLDERCAIINFRSS-QEWGWNDIHCHVPH 648

SEQ ID NO.

LS P G+RHWQWVD TPYNE+ TFWH EP++ +ERC ++NFR S + WGWND++C C-Type Lectin Receptor-Like:

Dendritic Cell Immunoreceptor: 167 LSDPEGQRHWQWVDQTPYNESSTFWHPREPSDPNERCVVLNFRKSPKRWGWNDVNCLGPQ 226

649 KSICEMKKIYI 681

C-Type Lectin Receptor-Like:

SEQ ID NO. 4

+S+CEM KI++

Dendritic Cell Immunoreceptor: 227 RSVCEMMKIHL 237

FIG. 2

The true of the tr

BLASTP ALIGNMENT OF C-TYPE LECTIN RECEPTOR-LIKE WITH C-TYPE LECTIN DDB27

Score = 529 (186.2 bits), Expect = 4.8e-50, P = 4.8e-50

Identities = 93/188 (49\$), Positives = 130/188 (69\$), Frame = +1

SEQ ID NO. 4 C-Type Lectin Receptor-Like:

DDB27:

118 VVSILLLSVCFTVSSVVPHNFMYSKTVKRLSKLREYQQYHSSLTCVMEGKDIED--WSCC 291

++

MSCC

LIFFLLLAISFFIAFVI----FFQKYSQLLEKKTTKELVHTTLECVKKNMPVEETAWSCC 106 57

SEQ ID NO. 4 C-Type Lectin Receptor-Like:

292 PTPWTSFQSSCYFISTGMQSWTKSQKNCSVMGADLVVINTTEEHDFIIHNLKRNSSYFLG 471 PW SF S+CYFIST SW S+K+C+ MAL+VINT EE DFI NL+ S+YF+G

107 PKNWKSFSSNCYFISTESASWQDSEKDCARMEAHLLVINTQEEQDFIFQNLQEESAYFVG 166

DDB27:

C-Type Lectin Receptor-Like:

SEQ ID NO. 4

472 LSHPRGRRHWQWVDHTPYNENVTFWHSGEPNNLDERCAIINFRSS-QEWGWNDIHCHVPH 648 EP++ +ERC ++NFR S + WGWND++C LS P G+RHWQWVD TPYNE+ TEWH 167 LSDPEGQRHWQWVDQTPYNESSTFWHPREPSDPNERCVVLNFRKSPKRWGWNDVNCLGPQ 226

DDB27:

649 KSICEMKKIYI 681

+S+CEM KI++

SEQ ID NO. 4 C-Type Lectin Receptor-Like: 227 RSVCEMMKIHL 237

DDB27:

FTG 3

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CARBOHYDRATE RECOGNITION DOMAIN) LECTIN RECEPTOR-LIKE, SUPERFAMILY MEMBER 6 BLASTP ALIGNMENT OF C-TYPE (CALCIUM DEPENDENT,

Score = 448 (157.7 bits), Expect = 1.8e-41, P = 1.8e-41

Identities = 92/209 (44%), Positives = 130/209 (62%), Frame = +1

49 PEEEP-QD-REKGLWWFQLKVWSMAVVSILLLSVCFTVSSVVPHNFMYSKTVKRLSKLRE 222 C-Type Lectin Receptor-Like: SEQ ID NO.

YS+ ++ L + S+ ++ +LLL++ F V+ ++ U P E+P +D R+ 29 PREKPIRDLRKPGSP--SLLLTSLMLL-LLLLAITFLVAFIIYFQ-KYSQLLEEKKAAKN 84

Mouse C-Type:

223 YQQYHSSLICVMEGKDIED--WSCCPTPWTSFQSSCYFIST--GMQSWTKSQKNCSVMGA 390 C-Type Lectin Receptor-Like:

85 IM--HNELNCTKSVSPMEDKVWSCCPKDWRLFGSHCYLVPTVSSSASWNKSEENCSRMGA 142 SW KS++NCS MGA CY + Tഗ Ĺτι ⋈ WSCCP + ED H+ I C

Mouse C-Type:

SEQ ID NO. 4

391 DLVVINTTEEHDFIIHNLKRNSSYFLGLSHPRGRRHWQWVDHTPYNENVTFWHSGEPNNL 570 C-Type Lectin Receptor-Like:

G R WQWVD TPY E++TFWH+GEP++ T +++XE+GT LVVI + EE DFI 143 HLVVIQSQEEQDFITGILDTHAAYFIGL-WDTGHRQWQWVDQTPYEESITFWHNGEPSSG 201

Mouse C-Type:

SEQ ID NO. 4

SEQ ID NO. 4

Mouse C-Type:

571 DERCAIINFRSSQEWGWNDIHCHVPHKSICEMKKI 675 C-Type Lectin Receptor-Like:

WGWNDI C + KS+C+MKKI +E+CA I +R

202 NEKCATIIYRWKTGWGWNDISCSLKQKSVCQMKKI 236